Figure 1. Structures of common compaction agents

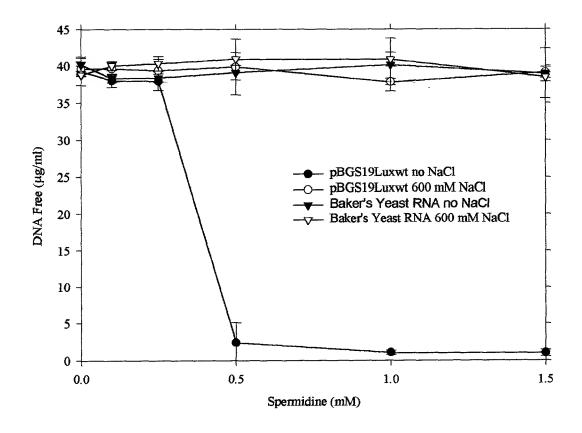


Figure 2. Precipitation by spermidine of  $40 \mu g/ml$  pBGS19Luxwt or Baker's yeast RNA in 10 mM Tris buffer at pH 8.0 with and without 600 mM NaCl. Error bars are  $\pm$ -one standard deviation.

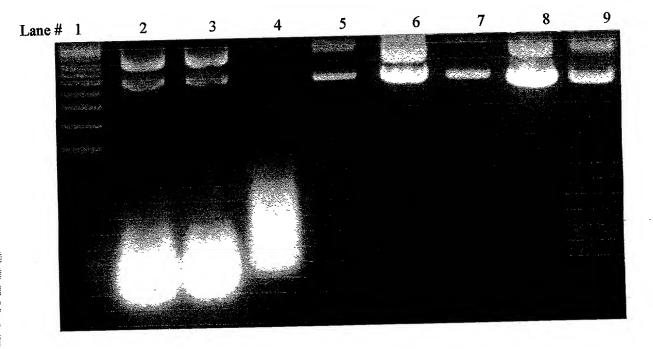


Figure 3. 1% agarose gel that traces the large-scale purification of pBGS19luxwt. Lane L is a supercoiled plasmid ladder from Gibco; Lane 2 is the preparation after Celite filtration, isopropanol precipitation, and resuspension; Lane 3 is the supernatant after LiCl precipitation; Lane 4 is the supernatant of the compaction precipitation; Lane 5 is the resuspended pellet of the compaction precipitation; Lane 6 is a 10X loading of the material in Lane 5; Lane 7 is after a Q sepharose anion exchange column (Fig. 5, bottom, Peak 5); Lane 8 is a 10X loading of Lane 7 and Lane 9 is pBGS19Luxwt plasmid DNA separated using the mini-prep procedure.

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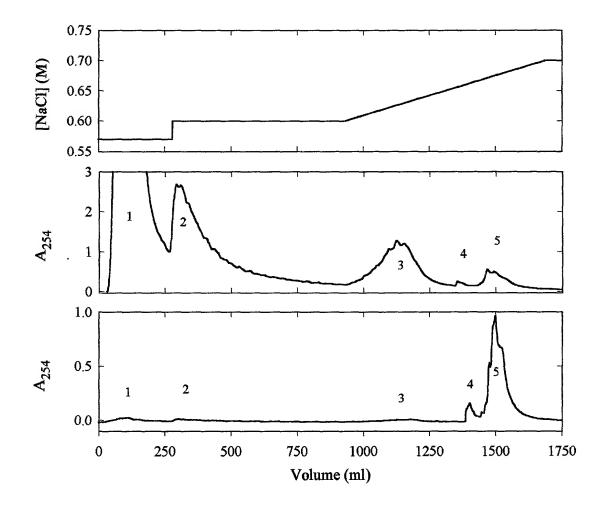


Figure 4. FPLC anion-exchange separation of pBGS19Luxwt of an alkaline lysate after isopropanol and LiCl precipitation. Top: NaCl gradient; Middle: with no previous compaction precipitation step; Bottom: identical separation after a previous compaction precipitation step.

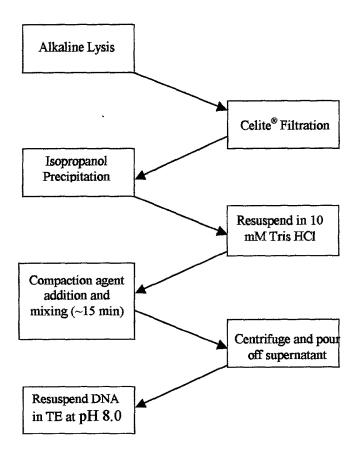


Figure 5. Summary of a selective precipitation-based noncolumn DNA purification as detailed in example 1.

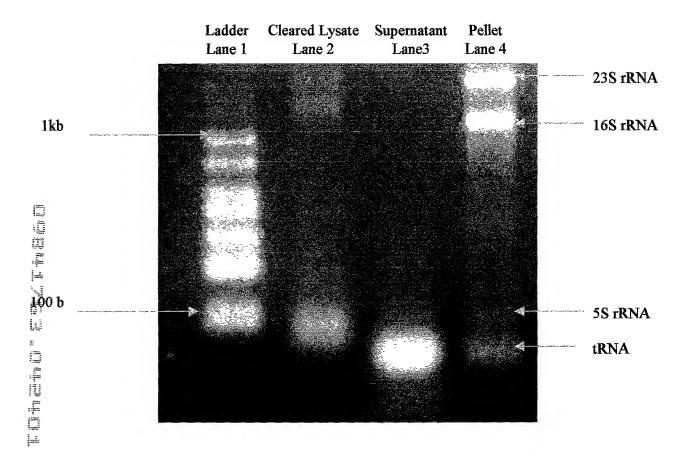


Figure 6. 3% biogel (from Bio101 Inc.) of V. proteolyticus RNA purified by Example 9. Lane 1 is the Ambion RNA Century Plus Size Markers; Lane 2 is the lysate after BPER addition, spermidine addition and centrifugation; Lane 3 is the supernatant of the 4 mM hexammine cobalt precipitation; and Lane 4 is the RNA pelleted in the hexammine cobalt precipitation but before any column separation.

1 [1]

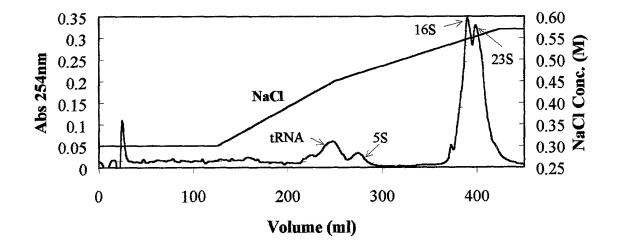


Figure 7. FPLC chromatogram of V. proteolyticus RNA on and 25 ml high performance Q Sepharose anion-exchange column (Pharmacia). The gradient was ran over 12 column volumes from 0.30 M NaCl to 0.57 M NaCl in a column buffer of 20 mM bis-tris propane and 20 mM EDTA at pH 6.9.

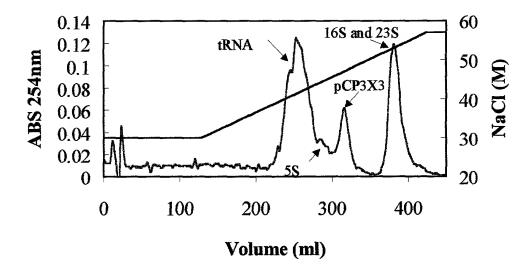


Figure 8. FPLC chromatogram of pCP3X3 aRNA containing *E. coli* strain JM109 on and 25 ml high performance Q Sepharose anion-exchange column (Pharmacia). The gradient was ran over 12 column volumes from 0.37 M NaCl to 0.57 M NaCl in a column buffer of 20 mM bis-tris propane and 20 mM EDTA at pH 6.9.

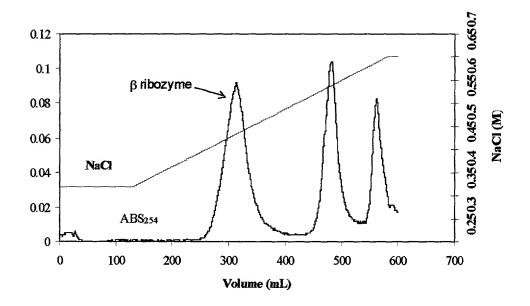
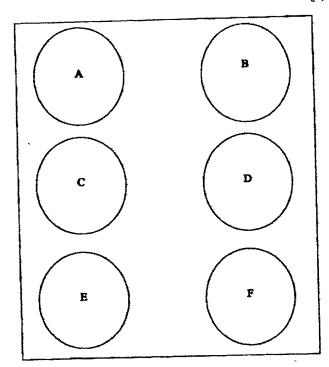


Figure 9. FPLC chromatogram of selective precipitation purified β ribozyme on and 25 ml high performance Q Sepharose anion-exchange column (Pharmacia). The gradient was ran over 12 column volumes from 0.37 M NaCl to 0.7 M NaCl in a column buffer of 10 mM bis-tris propane and 2 mM EDTA at pH 6.9.





## Description of Drawing ‡0B

## RNA isolation kit based on a hexammine cobalt precipitation process.

A. Lysia/DNA precipitation solution (e.g. 50% BPER with 2.5 mbd Spermidise or 1% Brij 38 with 2.5 mbd spermidise in 10 mbd bis tris proposes at pH 7). This may also be extended for use with plant cells and other colaryatio cells with the possibilities of homogenisation, ether lysis solutions, and breaking the lysis from the speculities DNA reasonal. Thus, there would be a lysis solution and a sequente DNA precipitation solution (e.g. 7 mbd becamenine colast for a total RNA precipitation of 4 mbd becamenine colast for a total RNA precipitation of 4 mbd becamenine colast for a total RNA precipitation of 4 mbd becamenine colast for a total RNA precipitation of 4 mbd becamenine colast for a total RNA precipitation of the first propose at pH 6.9 for a high molecular weight precipitated when solution B was used (e.g. 20 mbd hexamenine colast in 10 mbd bis tris propose at pH 6.9).

D. Sedaging solution (e.g. 50% inspropyl alcohol with 3 M Urea, 300 mbd NeCl, 25 mbd EDTA).

Total Robert weak.

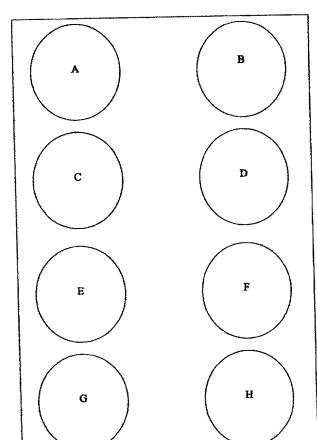
F. Final Robert weak.

Final Recompension colution (e.g. High pushly TE which is 10 and Tris and 1 mild EDTA at pH 2.0).

\*User may have to add enough alcohol to bring the solution to the proper percentage of alcohol.

Optional kit components that could be provided by and user:

Drawing 10/h 144



## Description of Drawing 10A

Pleanid DNA superation hit based on compaction agent procipitation technology. In the box a total of 8 solutions should be include two of which are optional. Solutions will include the 3 communa altafine lysis solutions, a low insis strength resuspension buffer, a compaction agent precipitation solution, and a chipleng solution, and optionally a 70% otheroil work solution and a final representation solution.

## Description of each bottle are as follows:

- A. Alkaline lysis solution I (e.g. 25 mM Tris and 10 mM EDTA at pH 8.0)

  B. Alkaline lysis solution II (e.g. 1% Sodium Dodocyl Sulliste (SDS) 0.2 N NgOH)

  C. Alkaline lysis solution III (e.g. 3 M KAc at pH 5.5)

  D. Resuperation solution (e.g. 10 mM Tris at pH 8.0)

  E. Compaction agent produktation solution (e.g. 2 mM Spermidine 3HCl and 10 mM Tris at pH 8.0)

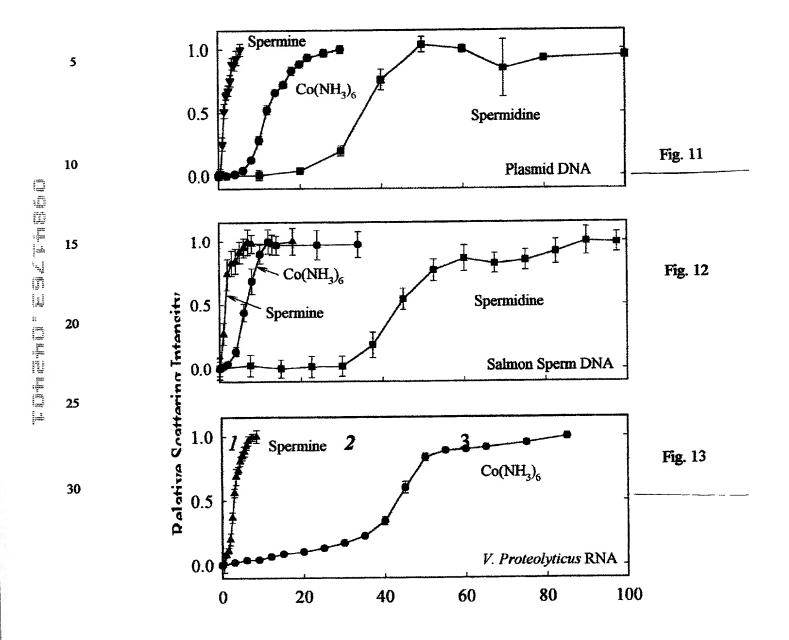
  F. Compaction agent produktation solution (e.g. 2 mM Spermidine 3HCl and 10 mM Tris at pH 8.0)

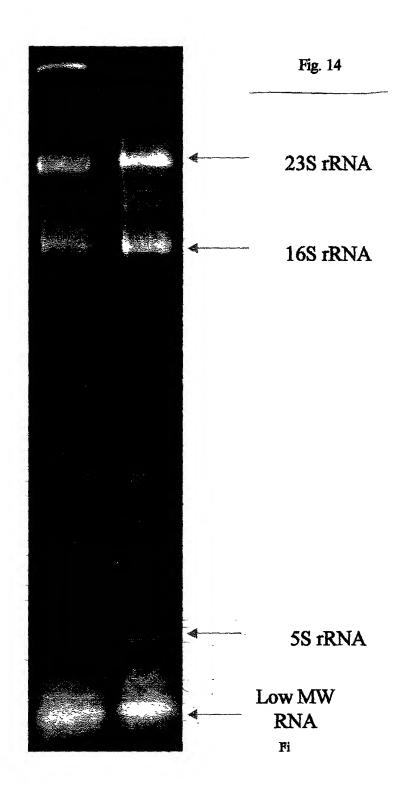
- F. Compaction agent stripping solution (e.g. 50% EtOH, 300 with NaCl, 12.5 mid EDTA') G. 70% EtOH wouth 's H. Flush Resuspension solution (e.g. High purity TE which is 10 mid This and 1 mid EDTA at pH 8.00."

"User may have to edd enough EtCHI to being the relation to the proper percentage of EIGH.

Optimal hit compensate that could be provided by end unor.

Also, solutions D and E can be combined to form a resespension/consequention agent precipitates solution.





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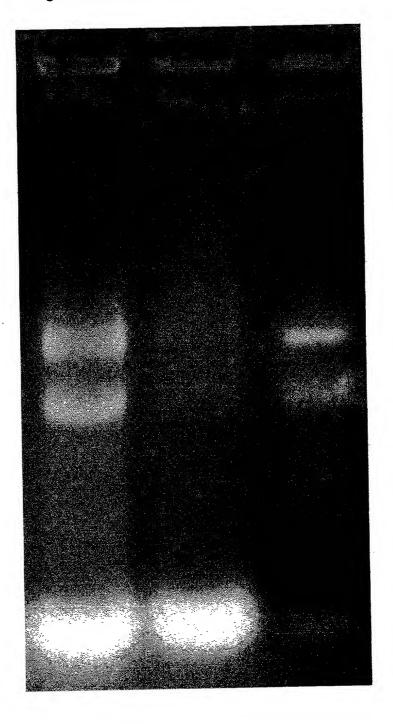
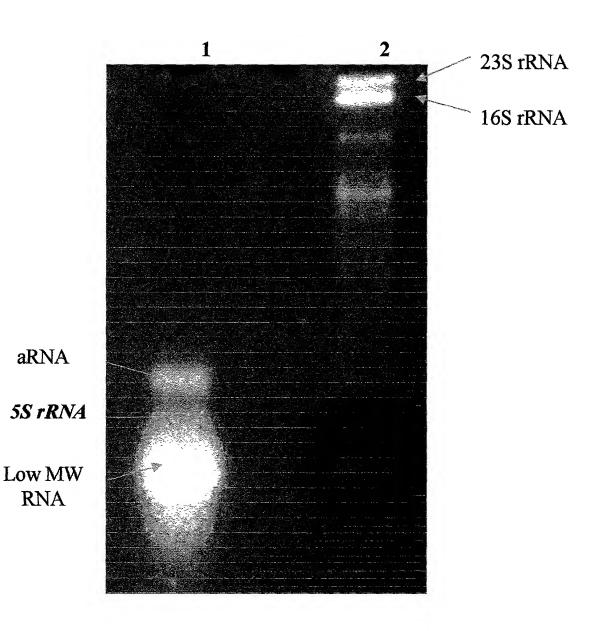


FIG. 15

FIG.16

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FIG.17

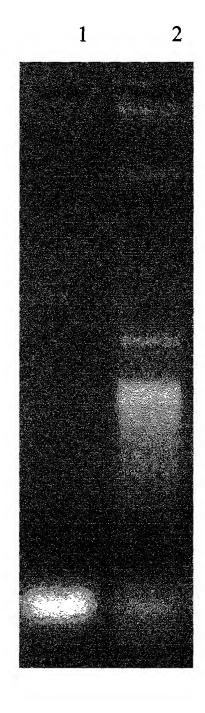


FIG.18

